

PCT

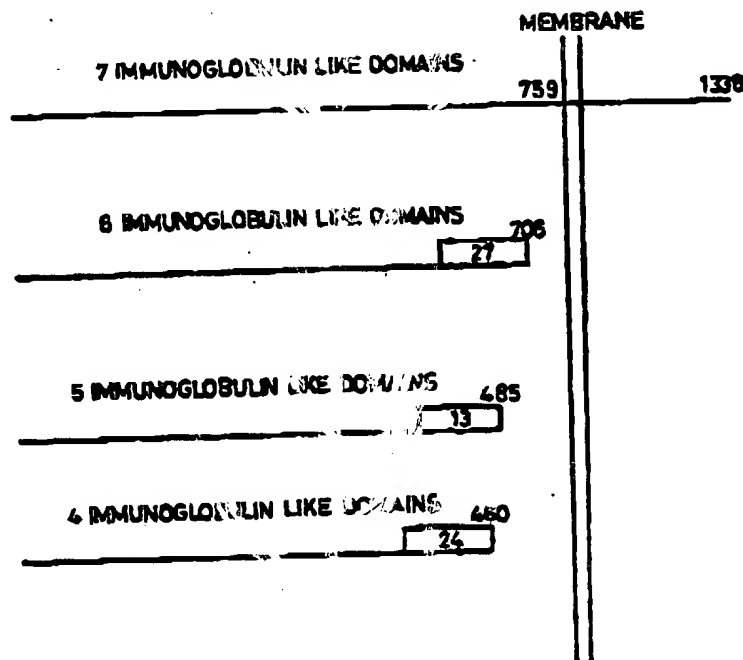
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(54) Title: FLT-1 (FMS-LIKE TYROSINE-KINASE), FLT-1 VARIANTS THEREOF USES AS GROWTH FACTOR INHIBITORS



(57) Abstract

Disclosed is an altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains, together with pharmaceutical compositions comprising the polypeptide, and various uses thereof.

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FLT-4 (fms-like Tyrosine kinase), FLT-15, variants thereof used as growth factor inhibitors

Field of the Invention

This invention relates to substances which inhibit growth factors, in particular, vascular endothelial growth factor (VEGF), methods of inhibiting growth factors and of treating tumours and regulating fertility.

Background of the Invention

A considerable number of human growth factors are now known, many of which have been at least partly characterised. Among them is vascular endothelial growth factor (VEGF), which has been identified in several tissues (Gospodarowicz *et al.*, 1989 PNAS 86, 7311-7315; Conn *et al.*, 1990 PNAS 87, 2628-2632; Tischer *et al.*, 1991 J. Biol. Chem. 266, 11947-11954). As its name suggests, this growth factor is a highly specific mitogen for endothelial cells and is greatly involved in angiogenesis. VEGF is a homodimeric glycoprotein of two 23kDa subunits exhibiting sequence homology with platelet-derived growth factor A and B chains and placenta growth factor.

The homologous tyrosine kinase receptors fms-like tyrosine kinase receptor (FLT) and kinase insert domain-containing receptor (KDR) function as high-affinity VEGF receptors (de Vries *et al.*, 1992 Science 255, 989-991; Terman *et al.*, 1992 Biochem. Biophys. Res. Commun. 187, 1579-1586). Both FLT and KDR are membrane-spanning receptors that each contain seven immunoglobulin-like domains in the extracellular ligand-binding region, an intracellular tyrosine kinase domain and a transmembrane domain. The transmembrane domain serves to anchor the receptor in the cell membrane of the cells in which it is expressed.

A number of membrane-bound receptor molecules have been found to exist in truncated soluble forms, generated either by proteolytic processing or by alternative splicing of

mRNA. Recently, Kendall & Thomas (1993 PNAS 90, 10,705-10,709, and WO94/21679) described the discovery of a soluble form of FLT receptor (sFLT) generated by alternative splicing.

Essentially, Kendall & Thomas screened a human umbilical vein endothelial cell (HUVEC) cDNA library with one probe specific for the 3' end of the flt coding region (encoding the intracellular tyrosine kinase domain) and with another probe specific for the 5' flt coding portion (encoding one of the extracellular N terminal domains). Clones which hybridised with the 5' specific probe but not with the 3' specific probe were selected for further study. In this way, a clone was isolated which encoded a soluble FLT polypeptide lacking the transmembrane domain and the intracellular domain. The truncation resulted from "readthrough" to an intronic termination codon. It was suggested by Kendall & Thomas that the soluble receptor could act as an efficient specific antagonist of VEGF *in vivo*.

The present invention is based on the discovery of further soluble variants of FLT, the existence of which was not predicted by the teaching of Kendall & Thomas.

Summary of the Invention

In a first aspect the invention provides an altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains. Preferably, the altered FLT polypeptide comprises four or fewer complete Ig-like domains. The altered soluble FLT polypeptide inhibits VEGF by preventing it binding to its natural receptors, flt and KDR, present on the surface of target cells. Surprisingly, such truncated forms, lacking a major extracellular portion of the molecule, are believed to retain affinity for VEGF.

The term "soluble" as used herein is intended to refer to altered forms of the FLT polypeptide which do not comprise a transmembrane domain and thus generally do not become associated with the cell membrane of cells in which the molecule is expressed. In particular, the invention provides soluble altered forms of the FLT polypeptide

consisting substantially of four or five complete immunoglobulin-like domains.

In a particular embodiment the invention provides an altered, soluble form of FLT having at its C-terminus a region substantially having the amino acid sequence of the sequences termed FLT4 or FLT15 shown in Figure 5, or a functional equivalent thereof. The term "functional equivalent" as used above is intended to include those polypeptides which have substantially the same deletions as the polypeptides encoded by FLT4 or FLT15 (with respect to the unaltered full length FLT molecule), but which may also have other deletions, additions or substitutions, (in particular conservative substitutions), and which retain an inhibitory effect for VEGF.

Preferably the polypeptide will also comprise, at its N-terminus, the amino acid sequence substantially corresponding to the equivalent portion of the unaltered wild-type FLT polypeptide. Conveniently, polypeptides in accordance with the invention will comprise around 400 to 500 amino acid residues, preferably around 480 amino acid residues, most preferably between 480 and 440 amino acid residues of the wild type FLT sequence.

Preferably the polypeptides of the invention arise by alternative splicing of mRNA or by proteolytic processing of a mature polypeptide, although it will be apparent to those skilled in the art that the polypeptide could be encoded by a nucleic acid derived, at least in part, by recombinant DNA technology.

In a further aspect the invention provides a nucleic acid sequence encoding a polypeptide in accordance with the invention. In a particular embodiment the invention provides a nucleic acid comprising the sequence of nucleotides inserted at position 1655 of the FLT 4 sequence shown in Figure 3 or the sequence of nucleotides inserted at position 1555 of the FLT 15 sequence shown in Figure 3, or a functional equivalent thereof. Examples of functionally equivalent nucleic acids include those sequences which encode substantially the same polypeptide as those encoded by FLT4 or FLT15 but which differ in nucleotide sequence as a result of the degeneracy of the genetic code. It will be apparent to those skilled in the art that the portion of the inserted nucleotide sequence in FLT4 and FLT15 occurring after the premature termination codon could be omitted without affecting the characteristics of the encoded polypeptide. Accordingly, nucleic acid molecules without

such sequences are also regarded as functionally equivalent for the purposes of the present invention.

Conveniently, the nucleic acid will substantially comprise the nucleotide sequence of FLT4 or FLT15 shown in Figure 3, together with the nucleotide sequence encoding the N-terminus of unaltered, wild-type FLT. Advantageously, the nucleic acid will be obtainable by means of PCR amplification from a sample of human cells. Desirably, the nucleic acid will be obtainable by means of PCR using primers intended to hybridise to non-conserved regions of the FLT molecule. Conveniently, the nucleic acid sequence will be obtainable by use of PCR primers designed to hybridise to the regions of the FLT sequence shown underlined in Figure 3, or immediately adjacent thereto. In particular, the PCR primers will conveniently have substantially the sequence: 5'- GCAAGGTGTGACTTTGTTC -3' and 5' - AGGATTCTTCCCCTGTGTA -3'.

In another aspect, the invention provides a method of inhibiting VEGF *in vitro*, comprising adding an effective amount of the polypeptide defined above. It may also be desirable to inhibit VEGF in a human subject. Thus the invention provides a method of inhibiting VEGF in a human subject, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. In particular, VEGF provides a mitogenic stimulus (particularly involved in angiogenesis), so inhibition of VEGF would be expected to provide therapeutic effects in the treatment of tumours or disorders involving inappropriate neovascularisation.

In particular the invention provides for a method of treating tumours or diseases involving inappropriate neovascularisation, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. Suitable diseases which might be amenable to treatment include ovarian cancer and ovarian hyperstimulation (Boocock *et al.*, 1995 *J. Natl. Cancer Inst.* 87, 506-516).

Furthermore, it has been conclusively demonstrated that FLT is expressed by trophoblasts and cells from ovarian and endometrial tissues (Charnock-Jones *et al.*, 1994 *Biology of Reproduction* 51, 524-530), which clearly suggests a role for VEGF in the growth and

differentiation of trophoblasts during implantation.

Thus, in particular, the invention provides a method of affecting the growth and/or migration of trophoblasts, ovarian or endometrial cells by inhibiting the action of VEGF, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance.

It will be appreciated by those skilled in the art that the identification of FLT on the surface of trophoblasts and endometrial cells also provides a number of possible methods of regulating fertility. For example, the growth of trophoblasts is essential for successful implantation of the embryo. Inhibition of trophoblast growth thus provides a method of contraception or contragestion.

Thus in a further aspect the invention provides a method of regulating the fertility of a human female, comprising administering an effective amount of the polypeptide defined above, together with a physiologically acceptable carrier substance. An "effective amount" of the polypeptide is an amount sufficient to substantially block the stimulus of VEGF on trophoblasts and/or endometrial cells. Typically, the method will result in reducing the fertility of the female.

Moreover, it might be possible to identify agents which can enhance the effect of VEGF on trophoblasts, and thereby improve the probability of successful implantation, either in assisted or spontaneous cycles. Candidates for such VEGF-enhancing agents would include anti-sense equivalents of the nucleic acid sequences encoding the truncated FLT polypeptides of the invention. It will be apparent to those skilled in the art that these could be used to improve the fertility of a human female.

In a further aspect the invention provides a pharmaceutical composition comprising the polypeptide defined above, together with a physiologically acceptable carrier substance. The composition could be used in vivo any one of the methods defined above. In yet another aspect the invention provides for the use of a polypeptide in accordance with the invention in the preparation of a therapeutic composition for the treatment of tumours and

diseases involving inappropriate neovascularisation. Examples of such conditions and diseases are detailed, inter alia, in WO94/10202 and WO94/21679. The invention also includes within its scope a method of making a pharmaceutical composition, comprising mixing the polypeptide defined above together with a physiologically acceptable carrier substance.

The invention will now be described by way of the following illustrative examples and with reference to the drawings, of which:

Figure 1 shows an amino acid multiple alignment of closely related tyrosine kinase receptors (flt, fms and kit, "kit" being another name for KDR);

Figure 2 shows typical results of agarose gel electrophoresis demonstrating the existence of alternatively-spliced flt-coding sequences in various tissue samples;

Figure 3 shows the nucleotide sequence of the 3' region of the sequences encoding full length VEGF receptors (FLT and the related receptor KDR), together with two sequences, FLT4 and FLT15, which encode polypeptides according to the invention;

Figure 4 is a schematic representation of wild type and mutant FLT molecules; and

Figure 5 shows the C terminal amino acid sequences of two polypeptides in accordance with the invention.

Example

Expression of FLT, the VEGF receptor, was investigated in cell lines derived from human trophoblast-like and ovarian and endometrial carcinomas. The trophoblast-like (choriocarcinoma) cell line used was BeWo (obtained from the American Type Culture Collection, Rockville MD, USA). The endometrial carcinoma cell lines were Ishikawa (obtained from Professor M Nishide, University of Tsukuba, Japan), and HEC 1-A and HEC 1-B (from ATCC, USA). The ovarian cancer cell lines were 7, 17R, 25, 25R and 35. These were all shown to be of epithelial origin and had been established in culture

for 10-30 passages. Cell lines 17R and 25R were derived after chemotherapy and subsequent relapse (line 25R originating from the same patient as line 25).

BeWo cells were grown in Ham's F12, according to ATCC recommendations. Endometrial carcinoma lines were grown in McCoy's medium (ICN Flow Laboratories, Irvine, UK) with 10% foetal calf serum (ICN Flow) plus 2mM L-glutamine (ICN Flow) and 50U/ml and 50mg/ml penicillin/streptomycin (ICN Flow).

It was decided to investigate expression of FLT in these cell lines and normal tissues by performing PCT and *in situ* hybridization. It was therefore necessary to construct suitable oligonucleotide primers and probes.

To help design appropriate primers, a protein multiple alignment of closely related tyrosine kinase receptors (FLT, FMS and KIT) was constructed (shown in Figure 1) using the computer program "pileup". This revealed regions of divergent sequence among this family of receptors. The regions chosen for primer design are shown with double underlining in Figure 1. The following nested PCR primers were then synthesized based on these protein sequences:

- A) 5' GCAAGGTGTGACTTTTGTTC 3'
- B) 5' GCGCTCGAGAGCATCACTCAG 3'
- C) 5' GCGCGGCCGCGAGTAAAATCCA 3'
- D) 5' AGGATTTCTTCCCCTGTGTA 3'

The underlined portions of these oligonucleotides are the regions which hybridise to the flt cDNA sequence. The other nucleotides were added to facilitate directional cloning. The cycles used were: first round with primers A and D [95°C 30 seconds, 55°C 30 seconds, 72°C 30 seconds] x 25; second round with primers B and C: [95°C 30 seconds, 44°C 30 seconds, 72°C 30 seconds] x 2 [95°C 30 seconds, 65°C 30 seconds, 72°C 30 seconds] x 25. The internal primers B and C had sites for the restriction enzymes Xho I and Eag I respectively at their 5' ends to permit directional cloning of the products.

It was found that certain tissues gave rise to PCR amplification products of notably larger size (as judged by agarose gel electrophoresis) than observed for the full length FLT cDNA product. Typical results are shown in Figure 2.

PCR products obtained using the nested set of primers A-D were run out on a gel. Lanes 1-3 are products obtained from primary tissue samples of the ovarian carcinomas designated 17, 17R and 25R. Lanes 4 to 7 are products obtained from cell lines established from the ovarian carcinomas 7, 17R, 25 and 25R. Lanes 8 to 10 are the cell lines HEC 1-A, HEC 1-B and Ishikawa respectively. Lane 11 contains products from HUVECs.

The standard size band was of the expected size (around 285bp) and was found to be identical to the 3' end of the published flt sequence (Shibuya *et al.*, 1990 *Oncogene* 5, 519-524). However it can be clearly seen that in addition to the full length flt cDNA PCR-amplified product, in lanes 2 (17R, primary tissue) and 4 (7, cell line) are larger bands of approximately 360bp. A faint band of similar size was also apparent in lane 5 (17R, cell line) but is not clearly seen when the gel photograph is reproduced. These larger bands were extracted from the gel by known techniques and subcloned into the plasmid vector pBluescript II KS and then subjected to sequence analysis using the dideoxynucleotide sequencing method (Sanger *et al.*, 1977 *PNAS* 71, 5463-5467).

Sequencing of five independent clones (Boocock *et al.*, 1995 *J. Natl. Cancer Inst.* 87, 506-516) revealed that each contained one of two novel insertions within the published flt sequence, in the region between the primers. Three of these clones (termed FLT5, FLT15 and FLT16) contained an 85bp insertion at about position 1555, whilst two other clones (FLT13 & FLT14) contained a 65bp insertion at about position 1665 (see Figure 3, numbering based on that of Shibuya *et al.*, 1990 cited above). The insertions account for the larger band size of the PCR products. However, both insertions contain an in-frame termination codon, so that corresponding full length RNAs would encode soluble, truncated receptor variants comprising the first five immunoglobulin-like domains of the extracellular region, up to amino acid 517 or 553, with either 24 or 14 (of which 13 are additional) unrelated amino acids at the C-terminus.

Although these variant flt clones were derived from partial cDNAs encoding only amino acids 503 onward, PCR products of the sizes predicted for corresponding full length cDNAs were amplified from cDNA derived from HUVEC cells, human chorion and ovarian carcinoma cell line 7, using primers specific for each of the novel insertions together with a primer binding just 5' of the initiating ATG (data not shown).

Figure 4 is a schematic representation of various FLT receptor molecules. At the top, (a) shows the wild type, full length FLT receptor molecule, (b) represents the truncated version described by Kendall & Thomas, (c) represents the polypeptide encoded by FLT4 and (d) represents the polypeptide encoded by FLT15. The numerals at the right show the number of amino acids in the molecule and numerals in the boxes represent the number of amino acids present in the sFLT variants but not in the wild type molecule.

Figure 5 shows the predicted C terminal amino acid sequence of the polypeptides which would be encoded by "full length" FLT4 and FLT15 clones (i.e. clones which contained all the nucleotide sequence 5' of the primer site used to generate the actual clones). The last 14 amino acids of the FLT4 clone, and the last 24 amino acids of the FLT15 clone, are divergent from the wild type FLT sequence.

Claims

1. An altered, soluble form of the FLT polypeptide being capable of binding to VEGF and thereby exerting an inhibitory effect thereon, the polypeptide comprising five or fewer complete immunoglobulin-like domains.
2. A polypeptide according to claim 1, comprising four or fewer complete immunoglobulin-like domains.
3. A polypeptide according to claim 1 or 2, having at its C terminus substantially the amino acid sequence of FLT4 as shown in Figure 5, or a functional equivalent thereof.
4. A polypeptide according to claim 1 or 2, having at its C terminus substantially the amino acid sequence of FLT15 as shown in Figure 5, or a functional equivalent thereof.
5. A polypeptide according to any one of the preceding claims, comprising around 400 to 500 amino acid residues of the wild type FLT polypeptide.
6. A nucleic acid sequence encoding a polypeptide in accordance with any one of the preceding claims.
7. A nucleic acid sequence according to claim 6, comprising the sequence of the nucleotides inserted at position 1655 of the FLT4 sequence shown in Figure 3, or a functional equivalent thereof.
8. A nucleic acid sequence according to claim 6, comprising the sequence of the nucleotides inserted at position 1555 of the FLT15 sequence shown in Figure 3, or a functional equivalent thereof.
9. A method of inhibiting VEGF *in vitro*, comprising adding an effective amount of a polypeptide in accordance with any one of claims 1 to 5.

10. A method of inhibiting VEGF in a human subject, comprising administering an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
11. A method according to claim 10, comprising the use of a polypeptide in accordance with any one of claims 1 to 5 in the treatment of tumours or diseases involving inappropriate neovascularisation.
12. A method according to claim 11, for the treatment of ovarian cancer, ovarian hyperstimulation, or endometriosis.
13. A method of affecting the growth and/or migration of trophoblasts, ovarian or endometrial cells by inhibiting the action of VEGF by administration of an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
14. A method of regulating the fertility of a human female by administration of an effective amount of a polypeptide in accordance with any one of claims 1 to 5, together with a physiologically acceptable carrier substance.
15. A pharmaceutical composition for use in the method of any one of claims 11 to 14, comprising a polypeptide in accordance with any one of claims 1 to 5, and a physiologically acceptable carrier substance.
16. A method of making a composition according to claim 15, comprising mixing a physiologically acceptable carrier substance together with a polypeptide according to any one of claims 1 to 5.

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1 50
 kit MRGARGAWDF LCVLLLLLLRV QTGSSQPSVS PGEPSPPSIH PGKSDLVVRV
 fmsM GPGVLLLLLV ATAWHGQIP VIEPSVP... ELVVKP
 flt MVSYWDTGVL LCALLSCLLL TGSSSGSKLK DPESLKG... TOHIMOA

51 100
 kit GDEIRLLCTD PGEVKW.... TFEILD ETNENKQ... NEWITE
 fms GATVTLRCVG NGSVENOGP. ASPHWT LYSDGSS... SILSTN
 flt GQTLHLQCRG EAAHKWSLPE MVSKESERLS ITKSACGRNG KQFCSTLTIN

101 150
 kit KAEATNTGKY TC....T... NKHGLSNSIY VFVRDPKLF... LVDRS
 fms NATFQNTGTY RC....TEPG DFI GGSAATH LYVKOPARFW... NVLAQE
 flt TAQANHTGFY SCKYLAVPTS KKETESAIY IFISDTGRPF VEMYSEIPEI

151 200
 kit LYGKEDNDTL VRCPLTDPEV .TNYSLKGCQ GKPLPKD.LR FIPDPKAGIM
 fms VVFEEDQDAL LPCLLTDPVL EAGVSLVRVR GRPLMRH.TN YSESPWHGFT
 flt IHMTIEGRELV IPCRVTSPII ..TVTLKKFP LDTLIPDGKR IIWDSRKGFI

201 250
 kit IKSVMKRAYHR LCLHCSVDOE GKSVLSEKFI LKVRPAFKAV PVVSUSKASY
 fms IHRAK.FIQS QDYQCSALMG GRKVMISISIR LKVQKVIQGP PALTLEFEL
 flt ISNAT.YKEI GLLTCEATVN GHLKYKINYL HRQNTIIDV QISTPRVKL

251 300
 kit LLREGEEFTV TCTI.KDVSS SVYSTWKREN SQTQLOEK... YNSWHHGD
 fms VRIRGEAAQI VCSA.SSVDV NEDVLOHNN ..TKLAIP... QOSDFHNN
 flt L..RGHTLV L NCTATPLNT RVQMTWSYPD EKNKRASVRR RIDQSNHAN

301 350
 kit FNYERQATLT ISSARVNDG VEMCYANNTF GSANVTITLE VVDKGFINI.
 fms .RYQKVLTLN LDQVDFQHAG NYSCVASNVQ GKHSTSMEEF VVESAYLNL.
 flt IFYS...VLT IDKMQNKDKG LYTCRVRSQP SFKSVNTSVH IYDKAFITVK

351 400
 kit FPMINTTVFV NDGENVDLIV EYEAFPKPEH QOWIYMNRTF TDKWEDYPKS
 fms SSEQNLIQEV TVGEGNLKLV MVEAYPGLOGFNWY LGPFSDHOPE
 flt HRKQOVLETV AGKRSYRLSM KVKAAPSPEV V..... WIKD

401 450
 kit ENESN..... IRYVSELHL TRKGTGGT YTEFVS..NS DVNAIAFNV
 fms PKLANATTKO TYRHTFTLSL PRKPSSEAGR YSEFAR..NP GGRALTFEL
 flt GLPATEKSAR YLTRGYSLLI KDVTEEDAGN YTILLSIKQS NVEKSLYATL

Fig. 1 Sheet 1

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	451		500
kit	YVNTKPEI..	LTYDRL....	...VN..GML QCVAAGPEP TIDWYFCPGT
fms	TLRYPPEV..	SVIWF....	...INGSGTL LCAASGYPOP NVTWLQCSGH
flt	IVNVKPOIYE	KAVSSFPDPA	LYPLGSRQIL TCTAYGIPOP TIKWFWHPCN
	501		550
kit	EQRC.....		
fms	TDRCD.....		
flt	HNHSEARCDF	CSNNEESFIL	LADSNMGNRI ESITORMAI EGKNKMASTL
	551		600
kitSASV
fmsEADV
flt	VVADSRISGI	YICIASNKVG	TVGRNISFYI TOVPNGEHVN LEKMPTEAD
	601		650
kit	LPV..DVQTL	NSSGPPF...	...GKLVVQSS
fms	LQVWDDPYPE	VLSQEPF...	...HKVTVQSL
flt	LKLSCTVNKE	LYROVTWILL	RTVMNRIMHY SISKOKMAIT KEHSITLNGT
	651		700
kit	IDSSAFKHNG	TVECKAYNDV G.....	
fms	LIVETLEHNQ	TYECRAHNSV G.....	
flt	IMNVSLQDSG	TYACRAPNVY	TGEEILOKKE ITIRDQEPY LLRNLSDEIV
	701		750
kit	..KTSAYFNE A.....	..FKGNKEQ	IHPHTLETP.....
fms	..SGSWAF.I P.....	..ISAGANTH	PPDEFLETP.....
flt	AISSSTTLDC	HANGVPEPQI	TWFKNNHKIQ QEPGIILGPG SSTLEIEFVT
	751		800
kitLLI GEVTVAGM/C
fmsVVV ACMSIMALL
flt	EEDEGVYHCK	ATNOKGSVES	SAYIVVQTS DKSNELELITL TCTC/ARTIF
	801		850
kit	IIVMILTYKY	LQKPMYEVOW	KVVEEINGNN YVYID..PTQ LPYDH.KWEE
fms	LLLLLLLYKY	KOKPKYQVW	KIIESYEGNS YTFID..PTQ LPYNE.KWEE
flt	WLLLTLLIRK	MKRSSEIKT	DYLSIMDPD EVPLDEQCR LPYDASKWEE
	851		900
kit	PRNRLSFGKT	LGAGAFGKV	EATAYGLIKS DAAMTVAVKM LKPSAHTER
fms	PRNNLOFGKT	LGAGAFGKV	EATAFGLGKE DAVLKVAVKM LKSTAHAEK
flt	ARERLKLKGS	LGRGAFGKV	QASAFGIKKS PTCRTVAVKM LKEGATASEY

Fig. 1 Sheet 2

SUBSTITUTE SHEET (RULE 26)

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	901		950
kit	EALMSELKVL SYLGNHMNIV NLLGACT.IG	GPTLVITEYC	CYGDLNFI R
fms	EALMSELKIM SHLGQHENIV NLLGACT.HG	GPVLVITEYC	CYGDLNFI R
flt	KALMTELKIL THIGHHLNVV NLLGACTKQG	GPLMVTVEYC	KYGILSNYIK
	951		1000
kit	RKRODFI... ..C...SKOE DHAEAAALYKN	L.....LHS	KESSCSDSTN
fms	RKAEAML... ..GPSLSPGQ DPEGGVYDYN	IHLEKKYVR	DSGESSQGV
flt	SKROLFFLNK DAALHMEPKK EKMEPGLEQG	KKRLDSVTS	SESFASSGQ
	1001		1050
kit	EXMDMKPGVS YVPTKADKR RSVRKGSYIE	ROVTPAIME	DELAIDLEDL
fms	TYVEMRP... ..VSTSSN... ..DSFSE	QOLD....KE	DGRPLELRL
flt	EDKSL..... ..SDVEE	EEDSDGFYKE	...PITMEDL
	1051		1100
kit	LSFSYQVAKG MAFLASKNCI HRDLAARNIL	LTHGRITKIC	DEGLARDIN
fms	LHFSYQVAKG MAFLASKNCI HRDVAARNVL	LTNGHVAKIG	DEGLARDIN
flt	ISYSFQVARG MEFLSSRKCI HRDLAARNIL	LSENWVKIC	DEGLARDIN
	1101		1150
kit	DSNYVVKGNA RLPVKWMAPE SIFNCVYTFE	SDVWSYGIFL	WEIFSLGSSP
fms	DSNYVVKGNA RLPVKWMAPE SIFDCVYTVQ	SDVWSYGILL	WEIFSLGLNP
flt	NPDYVRKGD T RLPVKWMAPE SIFDKIYSTK	SDVWSYGVLL	WEIFSLGGS?
	1151		1200
kit	YPGMPVDSKF YKMIKEGFRM LSPENAPAEM	YDIMKTOWDA	DPLKSTINQ
fms	YPGILVNSKF YKLVKDGYOM AOPAFAPKNI	YSIMQACWAL	EPTHRPTFQ
flt	YPGVQMDDEF CSRLREGMRM RAPEYSTPEI	YQIMLDQWHR	DPKERPREAE
	1201		1250
kit	IV....OLIE KQISES.TNH I.....Y	SNLANCSPNR	QKPVVDHSVR
fms	IC....SFLQ EQAQEDRRER D.....Y	TNLPSSRS.GGSGS
flt	LVEKLGDLLO ANVOQDGKDY IPINAILTGN	SGFTYSTPAF	SEDEFKESIS
	1251		1300
kit	INSVGSTASS SOP.....L LVHDDV....
fms	SSSELEESS SEH.....L TCEQGDIAQ	PLLOPNNYOF	C.....
flt	APKENSASSD DRYVNAFKE MSLEKIKTFE	ELLPNATSMF	DDYQGSSTL
	1301		1350
kit
fms
flt	LASPMLKFT WIDSKPKASL KIDLRTSKS	KESGLSDVSR	PSFCHSSCGH
	1351		1389
kit
fms
flt	VSEKGRFTY DHAELEKIA CCSPPPDYS	VVLYSTPPI	

Fig.1 Sheet 3

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1 2 3 4 5 6 7 8 9 10 11

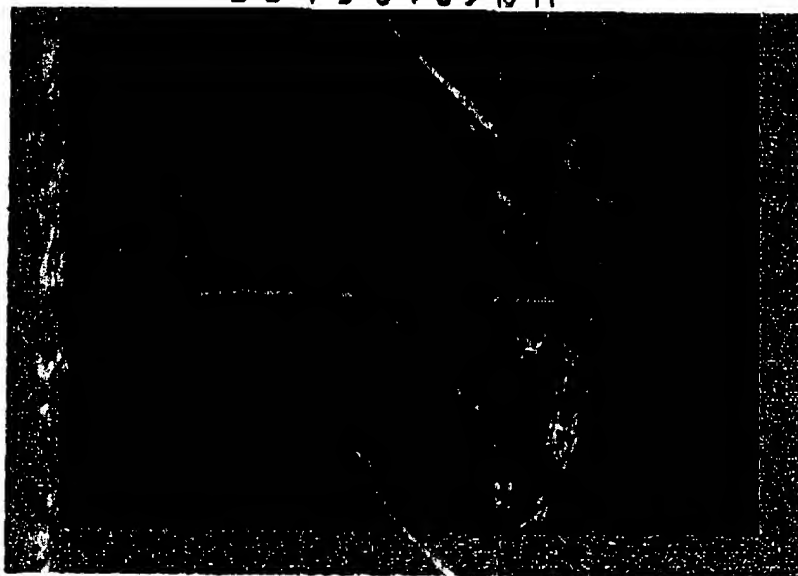


Fig. 2

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	1410				
KDR	AGAGTGC GCC	AACGAGCCCA	GCCAAGCTGT	CTCAGTGACA	AACCCATACC
FLT	ACCCCTGTAA	CCATAACATT	CCGAAGCAAG	GTGTGACTTT	TGTTCCAATA
	1460				
KDR	CTTGTGAAGA	ATGGAGAAGT	GTGGAGGACT	TCCAGGGAGG	AAATAAAATT
FLT	ATGAAGAGTC	CTTTATCCTG	GATGCTGACA	GCAACATGGG	AAACAGAATT
	1510				
KDR	GAAGTTAATA	AAAATCAATT	TGCTCTAATT	GAAGGAAAAA	ATAAA-----
FLT	GAGAGCATCA	CTCAGCGCAT	GGCAATAATA	GAAGGAAAGA	ATAAG-----
FLT4	GAGAGCATCA	CTCAGCGCAT	GGCAATAATA	GAAGGAAAGA	ATAAG-----
FLT15	GAGAGCATCA	CTCAGCGCAT	GGCAATAATA	GAAGGAAAGA	ATAAGCTTCC
KDR	-----	-----	-----	-----	-----
FLT	-----	-----	-----	-----	-----
FLT4	-----	-----	-----	-----	-----
FLT15	ACCAGCTGAC	AGTTCTTTCA	TGTTGCCACC	TACAAGCTTC	TGTTCCAACT
				1555	
KDR	-----	-----	-----	CTGTAAGTAC	CCTTGTTTATC
FLT	-----	-----	-----	ATGGCTAGCA	CCTTGGTTTGT
FLT4	-----	-----	-----	ATGGCTAGCA	CCTTGGTTTGT
FLT15	ACTTCCATTT	CCTTCCGTGA	CTCTAAACGG	ATGGCTAGCA	CCTTGGTTTGT
	1575				
KDR	CAAGCGGCAA	ATGTGTCAGC	TTTGTACAAA	TGTGAAGCGG	TCAACAAAGT
FLT	GGCTGACTCT	AGAATTTCTG	GAATCTACAT	TTGCATAGCT	TCCAATAAAG
FLT4	GGCTGACTCT	AGAATTTCTG	GAATCTACAT	TTGCATAGCT	TCCAATAAAG
FLT15	GGCTGACTCT	AGAATTTCTG	GAATCTACAT	TTGCATAGCT	TCCAATAAAG
	1625				
KDR	CGGGAGAGGA	GAGAGGGTGA	TCTCCTTCCA	CGTGACCAGG	-----
FLT	TTGGGACTGT	GGGAAGAAAC	ATAAGCTTTT	ATATCACAGA	-----
FLT4	TTGGGACTGT	GGGAAGAAAC	ATAAGCTTTT	ATATCACAGA	ATTGTCAAAC
FLT15	TTGGGACTGT	GGGAAGAAAC	ATAAGCTTTT	ATATCACAGA	-----
KDR	-----	-----	-----	-----	-----
FLT	-----	-----	-----	-----	-----
FLT4	TTTGAGTGCC	TTTATCCTTG	CTCTCAGGAA	TAGAAGCTCTA	CCTCATCGGA
FLT15	-----	-----	-----	-----	-----

Fig. 3 Sheet 1

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1665
 KDR -----GGTCC T---GAAATT ACTTTGCAAC CTGACATGCA GCCCACTGAG
 FLT -----TGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA TGCCGACGGA
 FLT4 TCTCATGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA TGCCGACGGA
 FLT15 -----TGTGC CAAATGGGTT TCATGTTAAC TTGGAAAAAA TGCCGACGGA

1710
 KDR CAGGAGAGCG TGTCTTTGTG GTGCACTGCA GACAGATCTA CGTTTGAGAA
 FLT AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC TTATACAGAG

FLT4 AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC TTATACAGAG
 FLT15 AGGAGAGGAC CTGAAACTGT CTTGCACAGT TAACAAGTTC TTATACAGAG

1760
 KDR CCTCACATGG TACAAGCTTG GCCCACAGCC TCTGCCAATC CAGGTGGGAG
 FLT ACGTTACTTG GATTTTACTG CGGACAGTTA ATAACAGAAC AATGCACTAC
 FLT4 ACGTTACTTG GATTTTACTG CGG
 FLT15 ACGTTACTTG GATTTTACTG CGG

1810
 FLT AGTATTAGCA AGCAAAAAAT GCCATCACT AAGGAGCACT CCTCACTCT

1860
 FLT TAATCTTACC ATCATGAAG TTTCCCTGCA AGATTCAGGC AATTATGCCT

1910
 FLT GCAGAGCCAG GAATGTATAC ACAGGGGAAG AAATCCTCCA GAAGAAAGAA

Fig. 3 Sheet 2

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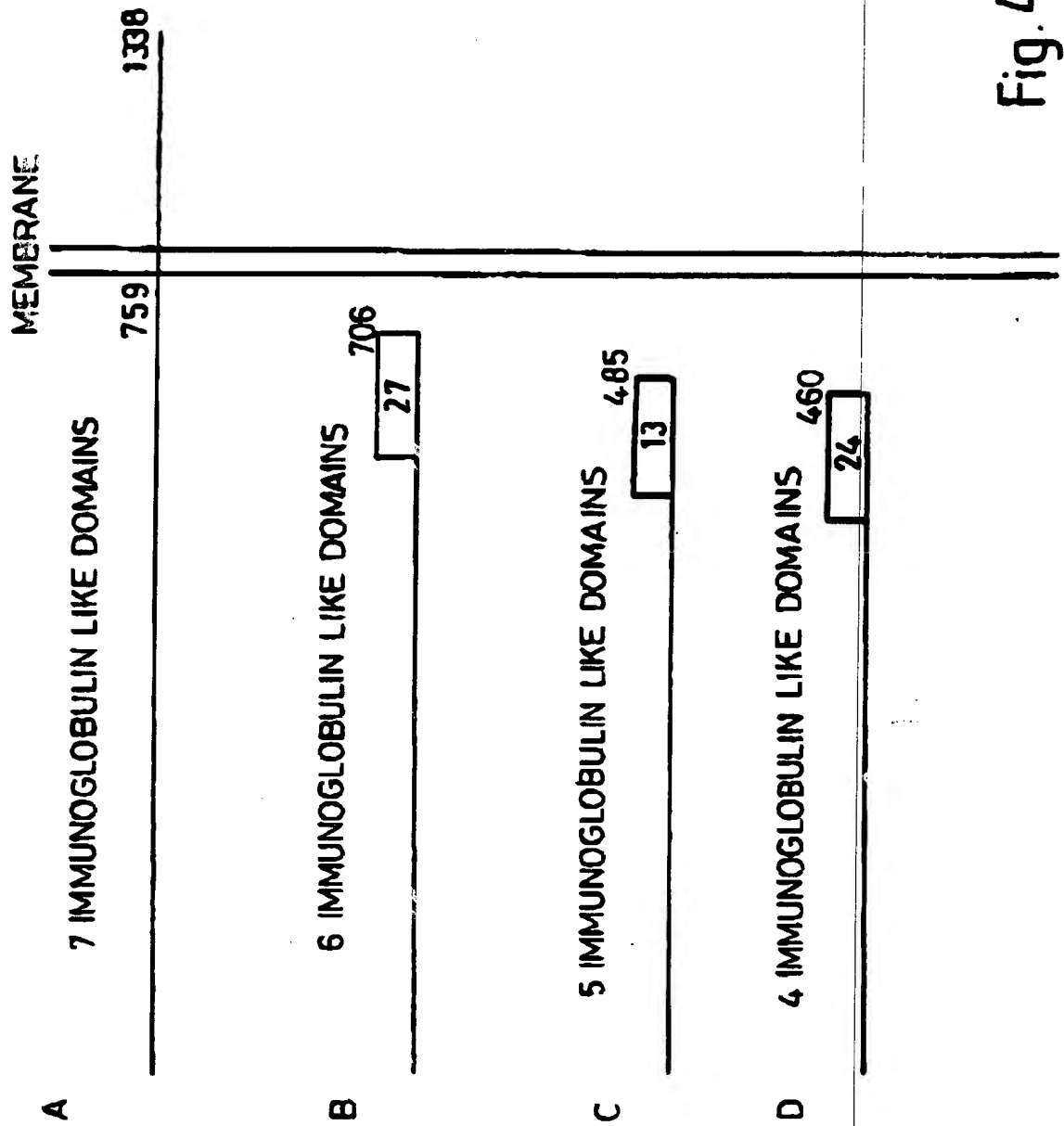


Fig. 4

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FLT4

1 ESITQRMAL EGKNKMASTL VVADSRISGI YICIASNKVG TVGRNISFYI

51 TELSNEFCLH PCSQE*

FLT15

1 ESITQRMAL EGKNKLPPAD SSFMLPPTSF SSNYFHFLP*

Fig. 5

INTERNATIONAL SEARCH REPORT

International Application No.
PCT/GB 95/01213

A. CLASSIFICATION OF SUBJECT MATTER
IPC 6 C12N15/12 C07K14/71 A61K38/17

According to International Patent Classification (IPC) or to both national classification and IPC

B. FIELDS SEARCHED

Minimum documentation searched (classification system followed by classification symbols)

IPC 6 C12N C07K

Documentation searched other than minimum documentation to the extent that such documents are included in the fields searched

Electronic data base consulted during the international search (name of data base and, where practical, search terms used)

C. DOCUMENTS CONSIDERED TO BE RELEVANT

Category *	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
------------	--	-----------------------

Y	CRIT REV ONCOG, 1993, 4 (6) P595-613, UNITED STATES, ROSNET O ET AL 'Hematopoietic receptors of class III receptor-type tyrosine kinases.' see the whole document	1,2,9-16
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Y	ONCOGENE, vol. 8, no. 11, November 1993 ENGLAND, pages 2931-2937, PAJUSOLA, K. ET AL.; 'Two human FLT4 receptor tyrosine kinase isoforms with distinct carboxy terminal tails are produced by alternative processing of primary transcripts' see the whole document	1,2,9-16
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☒ Further documents are listed in the continuation of box C.

☒ Patent family members are listed in annex.

* Special categories of cited documents:

- 'A' document defining the general state of the art which is not considered to be of particular relevance
- 'E' earlier document not published on or after the international filing date
- 'L' document which may throw doubts on priority claim(s) or which is cited to establish the publication date of another citation or other special reason (as specified)
- 'O' document referring to an oral disclosure, use, exhibition or other means
- 'P' document published prior to the international filing date but later than the priority date claimed

'T' later document published after the international filing date or priority date and not in conflict with the application but cited to understand the principle or theory underlying the invention

'X' document of particular relevance; the claimed invention cannot be considered novel or cannot be considered to involve an inventive step when the document is taken alone

'Y' document of particular relevance; the claimed invention cannot be considered to involve an inventive step when the document is combined with one or more other cited documents, such combination being obvious to a person skilled in the art.

'Z' document member of the same patent family

Date of the actual completion of the international search

17 October 1995

Date of mailing of the international search report

08. 11. 95

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Authorized officer

Nauche, S

INTERNATIONAL SEARCH REPORT

International Application No
PCT/GB 95/01213

C.(Continuation) DOCUMENTS CONSIDERED TO BE RELEVANT

Category	Citation of document, with indication, where appropriate, of the relevant passages	Relevant to claim No.
Y	ONCOGENE, AUG 1993, 8 (8) P2293-8, ENGLAND, FINNERTY H ET AL 'Molecular cloning of murine FLT and FLT4.' see the whole document ---	1,2,9-16
Y	WO,A,94 01576 (SYSTEMIX INC) 20 January 1994 see the whole document ---	1,2,9-16
A	WO,A,93 15201 (NEW ENGLAND DEACONESS HOSPITAL) 5 August 1993 see the whole document ---	1-16
A	WO,A,92 14748 (AMERICAN CYANAMID CO) 3 September 1992 see the whole document -----	1-16

INTERNATIONAL SEARCH REPORT

Intern. application No.

PCT/GB95/01213

Box I Observations where certain claims were found unsearchable (Continuation of item 1 of first sheet)

This international search report has not been established in respect of certain claims under Article 17(2)(a) for the following reasons:

1. ☒ Claims Nos.: 10-13
because they relate to subject matter not required to be searched by this Authority, namely:
Remark: Although claims 10-13 are directed to a method of treatment of the human/animal body as well as diagnostic methods (Rule 39.1(iv) PCT) the search has been carried out and based on the alleged effects of the compound/composition.
2. ☐ Claims Nos.:
because they relate to parts of the international application that do not comply with the prescribed requirements to such an extent that no meaningful international search can be carried out, specifically:
3. ☐ Claims Nos.:
because they are dependent claims and are not drafted in accordance with the second and third sentences of Rule 6.4(a).

Box II Observations where unity of invention is lacking (Continuation of item 2 of first sheet)

This International Searching Authority found multiple inventions in this international application, as follows:

1. ☐ As all required additional search fees were timely paid by the applicant, this international search report covers all searchable claims.
2. ☐ As all searchable claims could be searched without effort justifying an additional fee, this Authority did not invite payment of any additional fee.
3. ☐ As only some of the required additional search fees were timely paid by the applicant, this international search report covers only those claims for which fees were paid, specifically claims Nos.:
4. ☐ No required additional search fees were timely paid by the applicant. Consequently, this international search report is restricted to the invention first mentioned in the claims; it is covered by claims Nos.:

Report on Protest

- ☐ The additional search fees were accompanied by the applicant's protest.
- ☐ No protest accompanied the payment of additional search fees.

INTERNATIONAL SEARCH REPORT

International Application No.
PCT/GB 95/01213

Patent document cited in search report	Publication date	Patent family member(s)	Publication date
WO-A-9401576	20-01-94	AU-B- 4667593	31-01-94
		CA-A- 2135193	20-01-94
		EP-A- 0654088	24-05-95
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		CA-A- 2128722	05-08-93
		EP-A- 0624192	17-11-94
		JP-T- 7504813	01-06-95
WO-A-9214748	03-09-92	EP-A- 0536350	14-04-93